



Figure 1B

SEQ ID NO:26	* *** ** * ** ** * ** * ** * ** * ** *	SSTHLTPYRVVILRLIILGFFLQYRCRTHPVKDAYPLWLTSVICEVWFALSWLLDQFPKW	360
SEQ ID NO:30		ASSKINPYRMVIVLRLVLSIFLHYRLTNVPVNAYPLWLLSVICEIWFALSWLLDQFPKW	
SEQ ID NO:33		PSSRINPYRMVIMLRLVILCLFLHYRITNPVPNAFALWLVSVICEIWFALSWLLDQFPKW	361
SEQ ID NO:26	* ***** * ***** * ***** * ***** *	SPVNRETYLDRLSMRFDREGEPSQLAPIDVFVSTVDPLKEPPLVTANTVLSILAVDYPVD	420
SEQ ID NO:30		FPINRETYLDRALRYDREGEPSQLAAVDIFVSTVDPLKEPPIVTANTVLSILAVDYPVD	
SEQ ID NO:33		FPVNRETYLDRALRYDREGEPSQLAAVDIFVSTVDPLKEPPLVTANTVLSILAVDYPVD	421
SEQ ID NO:26	***** ** * ** ** ** ** ** ** ** ** ** ** *	KVSCYVSDDGASMLTFEALSETAEFAKKWAPFCCKHKSIEPRAPEFYFAQKIDYLDKDKVQP	480
SEQ ID NO:30		KVSCYVSDDGASMLTFDALAETSEFARKWVPFVKKYDIEPRAPEFYFCQKIDYLDKDKVQP	
SEQ ID NO:33		KVSCYVFDDGAAMLSFESLAETSEFARKWVPFCKKYSIEPRAPEWYFAAKIDYLDKDKVQT	481
SEQ ID NO:26	***** ***** ** * ***** * ***** * ***** *	SFVKERRAMKREYEEFKVIRINALVAKAQKVP EEGWMTMQDGTWP GNNNRDHPGMIQVFLG	540
SEQ ID NO:30		SFVKDRRAMKREYEEFKIRINALVSKALKVP EEGWIMQDGTWP GNNTRDHPGMIQVFLG	
SEQ ID NO:33		SFVKDRRAMKREYEEFKIRINALVSKALKCP EEGWVMQDGTWP GNNTGDPGMIQVFLG	541
SEQ ID NO:26	** * ***** ***** ***** ***** ***** ** *****	HSGGFDTEGNELPRLVYVSREKRP GFQHHKKAGAMNALIRVSAVL TN GAYLLNVDCDHYF	600
SEQ ID NO:30		HSGGLDTEGNELPRLVYVSREKRP GFQHHKKAGAMNALIRVSAVL TN GQYMLNLDCDHYI	
SEQ ID NO:33		QNGGLDAEGNELPRLVYVSREKRP GFQHHKKAGAMNALIRVSAVL TN GPFILNLDCDHYI	601



Figure 1D

SEQ ID NO:26	*** ** * * * * * ** * * * * * ** * * * * * ** * * * * *	901
SEQ ID NO:30	PLLAYCTLPAICLLTGKFIVPEISNYASIWFILLFVSIFSTGILELRWSGVLTLEDWRNE	
SEQ ID NO:33	PLLVCILPAICLLTGKFIMPEISNLSIWFIALFLSIFATGILEMRWSGVGIDEWWRNE	
	PLLMYCTLLAVCLFTNQFIIPQISNIAISIWFSLFLSIFATGILEMRWSGVGIDEWWRNE	960
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SEQ ID NO:26	QFWVIGGTSAHLFAVFQGLLKVLGIDTNTFTVTSKASDEDDGFAELYVFKWTSLLIPPTT	
SEQ ID NO:30	QFWVIGGISAHLFAVFQGLLKVLGIDTNTFTVTSKANDEEGDFAELYMFKWTLLIPPTT	
SEQ ID NO:33	QFWVIGGVS AHLFAVFQGLKVLGIDTNTFTVTSKASDEDDGFAELYLFKWTLLIPPTT	1020
	***** ** * * * * * ** * * * * * ** * * * * * ** * * * * *	
SEQ ID NO:26	ILVNMVGIVAGVSFAINSYGYSWGPLFGRLFFAIWVIVHLYPFLKGLLGRQNRTPPTVI	
SEQ ID NO:30	ILIINMVGIVAGTSYAINSYGYSWGPLFGKLFFAFWVIVHLYPFLKGLMGRQNRTPPTVI	
SEQ ID NO:33	LLIVNLVGVVAGVSYAINSYGYSWGPLFGKLFFAFWVIVHLYPFLKGLMGRQNRTPPTIV	1080
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SEQ ID NO:26	VWSVLLASIFSLWVRIDPFTSDSTKAR-GQCGIDC	
SEQ ID NO:30	VWAVLLASIFSLWVRVDPFTTTRLAGPNIQTGGINC	
SEQ ID NO:33	VWSVLLASIFSLWVRIDPFTSRVTGPDILECGINC	1116
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	***** ** * * * * * ** * * * * * ** * * * * * ** * * * * *	1081